

#6

SEQUENCE LISTING



<110> RIKEN

AJINOMOTO CO., INC.

<120> A METHOD FOR INCREASING STRESS-RESISTANCE TO A PLANT

<130> 204936US-3524-10-0

<150> JP 2001-72650

<151> 2001-03-14

<160> 15

<170> PatentIn Ver. 2.0

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<211> 344

<212> PRT

<213> Arabidopsis thaliana

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25

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Val Thr Phe Leu Ala Gly Asn Gly Asp Tyr Val Lys Gly Val Val Gly

35

40

45

Leu Ala Lys Gly Leu Arg Lys Val Lys Ser Ala Tyr Pro Leu Val Val

50

55

60

Ala Met Leu Pro Asp Val Pro Glu Glu His Arg Arg Ile Leu Val Asp

65

70

75

80

Gln Gly Cys Ile Val Arg Glu Ile Glu Pro Val Tyr Pro Pro Glu Asn

85

90

95

Gln Thr Gln Phe Ala Met Ala Tyr Tyr Val Ile Asn Tyr Ser Lys Leu

100

105

110

Arg Ile Trp Lys Phe Val Glu Tyr Ser Lys Met Ile Tyr Leu Asp Gly

115

120

125

Asp Ile Gln Val Tyr Glu Asn Ile Asp His Leu Phe Asp Leu Pro Asp

130

135

140

Gly Tyr Leu Tyr Ala Val Met Asp Cys Phe Cys Glu Lys Thr Trp Ser

145

150

155

160

His Thr Pro Gln Tyr Lys Ile Arg Tyr Cys Gln Gln Cys Pro Asp Lys

165

170

175

Val Gln Trp Pro Lys Ala Glu Leu Gly Glu Pro Pro Ala Leu Tyr Phe

180

185

190

Asn Ala Gly Met Phe Leu Tyr Glu Pro Asn Leu Glu Thr Tyr Glu Asp

195

200

205

Leu Leu Arg Thr Leu Lys Ile Thr Pro Pro Thr Pro Phe Ala Glu Gln

210

215

220

Asp Phe Leu Asn Met Tyr Phe Lys Lys Ile Tyr Lys Pro Ile Pro Leu

225

230

235

240

Val Tyr Asn Leu Val Leu Ala Met Leu Trp Arg His Pro Glu Asn Val

245

250

255

Glu Leu Gly Lys Val Lys Val Val His Tyr Cys Ala Ala Gly Ser Lys

260

265

270

Pro Trp Arg Tyr Thr Gly Lys Glu Ala Asn Met Glu Arg Glu Asp Ile

275

280

285

Lys Met Leu Val Lys Lys Trp Trp Asp Ile Tyr Asp Asp Glu Ser Leu

290

295

300

Asp Tyr Lys Lys Pro Val Thr Val Val Asp Thr Glu Val Asp Leu Val

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310

315

320

Asn Leu Lys Pro Phe Ile Thr Ala Leu Thr Glu Ala Gly Arg Leu Asn

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335

Tyr Val Thr Ala Pro Ser Ala Ala

340

<210> 2

<211> 335

<212> PRT

<213> Arabidopsis thaliana

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Met Ala Pro Glu Ile Asn Thr Lys Leu Thr Val Pro Val His Ser Ala

1 5 10 15

Thr Gly Gly Glu Lys Arg Ala Tyr Val Thr Phe Leu Ala Gly Thr Gly

20 25 30

Asp Tyr Val Lys Gly Val Val Gly Leu Ala Lys Gly Leu Arg Lys Ala

35 40 45

Lys Ser Lys Tyr Pro Leu Val Val Ala Val Leu Pro Asp Val Pro Glu

50 55 60

Asp His Arg Lys Gln Leu Val Asp Gln Gly Cys Val Val Lys Glu Ile

65 70 75 80

Glu Pro Val Tyr Pro Pro Glu Asn Gln Thr Glu Phe Ala Met Ala Tyr

85 90 95

Tyr Val Ile Asn Tyr Ser Lys Leu Arg Ile Trp Glu Phe Val Glu Tyr

100 105 110

Asn Lys Met Ile Tyr Leu Asp Gly Asp Ile Gln Val Phe Asp Asn Ile

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125

Asp His Leu Phe Asp Leu Pro Asn Gly Gln Phe Tyr Ala Val Met Asp

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135

140

Cys Phe Cys Glu Lys Thr Trp Ser His Ser Pro Gln Tyr Lys Ile Gly

145

150

155

160

Tyr Cys Gln Gln Cys Pro Asp Lys Val Thr Trp Pro Glu Ala Lys Leu

165

170

175

Gly Pro Lys Pro Pro Leu Tyr Phe Asn Ala Gly Met Phe Val Tyr Glu

180

185

190

Pro Asn Leu Ser Thr Tyr His Asn Leu Leu Glu Thr Val Lys Ile Val

195

200

205

Pro Pro Thr Leu Phe Ala Glu Gln Asp Phe Leu Asn Met Tyr Phe Lys

210

215

220

Asp Ile Tyr Lys Pro Ile Pro Pro Val Tyr Asn Leu Val Leu Ala Met

225

230

235

240

Leu Trp Arg His Pro Glu Asn Ile Glu Leu Asp Gln Val Lys Val Val

245

250

255

His Tyr Cys Ala Ala Gly Ala Lys Pro Trp Arg Phe Thr Gly Glu Glu

260

265

270

Glu Asn Met Asp Arg Glu Asp Ile Lys Met Leu Val Lys Lys Trp Trp

275

280

285

Asp Ile Tyr Asn Asp Glu Ser Leu Asp Tyr Lys Asn Val Val Ile Gly

290

295

300

Asp Ser His Lys Lys Gln Gln Thr Leu Gln Gln Phe Ile Glu Ala Leu

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335

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<212> DNA

<213> Arabidopsis thaliana

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gattacgtga aaggagtcgt tggtttagcc aaaggggtta ggaaagtcaa atcggcttat 180  
ccactcgtag tagcgatgtt acccgacgic ccggaggaac accgtcgtat acttgtggat 240  
caaggatgca tcgtccgtga aatcgaaccc gttaaccac ccgagaacca aactcagttc 300  
gccatggcctt attacgtcat caactactct aaactccgta tctggaagtt tgtggagtat 360  
agtaaaatga tatattaga tggagacatt caagtttacg aaaacatcga tcacttgttt 420  
gacctaccag atggctatit gtacgcggtg atggattgtt tctgtgagaa aacatggagt 480  
cacacgccgc aatacaagat cagatattgc caacaatgcc ccgacaaagt ccagtggcca 540  
aaagcggagc ttggagagcc accggctctt tacttcaacg ccggaatgtt cttgtacgag 600

cctaacctcg agacitacga ggatctacta cgaacactta aaatcactcc tccgactcct 660  
 ttcgctgaac aggatttttt gaacatgtac ttttaagaaaa tctacaagcc gattccttta 720  
 gtgtacaatc tcgtccttgc gatgttaagg cgtcaccagc aaaatgtaga gcttggaaaa 780  
 gtcaaggatg ttcactactg tgcagcgggt tcgaagccgt ggagatacac agggaaagaa 840  
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 gacgaatcct tggattacaa gaaacctgtt accgttgttg acacagaggt cgatctcgtg 960  
 aatctgaagc cgttcacac cgctcttact gaagctggcc ggctcaacta cgtgaccgca 1020  
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<213> Artificial Sequence

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caaggatccg cagatcacgt gctaatac

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<223> Description of Artificial Sequence:primer

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cgcggatccc tggigtgac aagaacctcg etc

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SEQUENCE LISTING

<110> TAJI, Teruaki  
SHINOZAKI, Kazuo  
OHSUMI, Chieko



<120> A METHOD FOR INCREASING STRESS-RESISTANCE TO A PLANT

<130> 20436US0

<140> 09/810,506

<141> 2001-03-19

<150> JP2001-072650

<151> 2001-03-14

<160> 15

<170> PatentIn version 3.1

<210> 1

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<212> PRT

<213> Arabidopsis thaliana

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Met Ala Pro Gly Leu Thr Gln Thr Ala Asp Ala Met Ser Thr Val Thr  
 1 5 10 15  
 Ile Thr Lys Pro Ser Leu Pro Ser Val Gln Asp Ser Asp Arg Ala Tyr  
 20 25 30  
 Val Thr Phe Leu Ala Gly Asn Gly Asp Tyr Val Lys Gly Val Val Gly  
 35 40 45  
 Leu Ala Lys Gly Leu Arg Lys Val Lys Ser Ala Tyr Pro Leu Val Val  
 50 55 60  
 Ala Met Leu Pro Asp Val Pro Glu Glu His Arg Arg Ile Leu Val Asp  
 65 70 75 80  
 Gln Gly Cys Ile Val Arg Glu Ile Glu Pro Val Tyr Pro Pro Glu Asn  
 85 90 95  
 Gln Thr Gln Phe Ala Met Ala Tyr Tyr Val Ile Asn Tyr Ser Lys Leu  
 100 105 110  
 Arg Ile Trp Lys Phe Val Glu Tyr Ser Lys Met Ile Tyr Leu Asp Gly  
 115 120 125  
 Asp Ile Gln Val Tyr Glu Asn Ile Asp His Leu Phe Asp Leu Pro Asp  
 130 135 140  
 Gly Tyr Leu Tyr Ala Val Met Asp Cys Phe Cys Glu Lys Thr Trp Ser  
 145 150 155 160  
 His Thr Pro Gln Tyr Lys Ile Arg Tyr Cys Gln Gln Cys Pro Asp Lys  
 165 170 175  
 Val Gln Trp Pro Lys Ala Glu Leu Gly Glu Pro Pro Ala Leu Tyr Phe  
 180 185 190  
 Asn Ala Gly Met Phe Leu Tyr Glu Pro Asn Leu Glu Thr Tyr Glu Asp  
 195 200 205

Leu Leu Arg Thr Leu Lys Ile Thr Pro Pro Thr Pro Phe Ala Glu Gln  
210 215 220

Asp Phe Leu Asn Met Tyr Phe Lys Lys Ile Tyr Lys Pro Ile Pro Leu  
225 230 235 240

Val Tyr Asn Leu Val Leu Ala Met Leu Trp Arg His Pro Glu Asn Val  
245 250 255

Glu Leu Gly Lys Val Lys Val Val His Tyr Cys Ala Ala Gly Ser Lys  
260 265 270

Pro Trp Arg Tyr Thr Gly Lys Glu Ala Asn Met Glu Arg Glu Asp Ile  
275 280 285

Lys Met Leu Val Lys Lys Trp Trp Asp Ile Tyr Asp Asp Glu Ser Leu  
290 295 300

Asp Tyr Lys Lys Pro Val Thr Val Val Asp Thr Glu Val Asp Leu Val  
305 310 315 320

Asn Leu Lys Pro Phe Ile Thr Ala Leu Thr Glu Ala Gly Arg Leu Asn  
325 330 335

Tyr Val Thr Ala Pro Ser Ala Ala  
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<210> 2

<211> 335

<212> PRT

<213> Arabidopsis thaliana

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Met Ala Pro Glu Ile Asn Thr Lys Leu Thr Val Pro Val His Ser Ala  
1 5 10 15

Thr Gly Gly Glu Lys Arg Ala Tyr Val Thr Phe Leu Ala Gly Thr Gly  
20 25 30

Asp Tyr Val Lys Gly Val Val Gly Leu Ala Lys Gly Leu Arg Lys Ala  
35 40 45

Lys Ser Lys Tyr Pro Leu Val Val Ala Val Leu Pro Asp Val Pro Glu  
50 55 60

Asp His Arg Lys Gln Leu Val Asp Gln Gly Cys Val Val Lys Glu Ile  
65 70 75 80

Glu Pro Val Tyr Pro Pro Glu Asn Gln Thr Glu Phe Ala Met Ala Tyr  
85 90 95

Tyr Val Ile Asn Tyr Ser Lys Leu Arg Ile Trp Glu Phe Val Glu Tyr  
100 105 110

Asn Lys Met Ile Tyr Leu Asp Gly Asp Ile Gln Val Phe Asp Asn Ile  
115 120 125

Asp His Leu Phe Asp Leu Pro Asn Gly Gln Phe Tyr Ala Val Met Asp  
130 135 140

Cys Phe Cys Glu Lys Thr Trp Ser His Ser Pro Gln Tyr Lys Ile Gly  
145 150 155 160

Tyr Cys Gln Gln Cys Pro Asp Lys Val Thr Trp Pro Glu Ala Lys Leu  
165 170 175

Gly Pro Lys Pro Pro Leu Tyr Phe Asn Ala Gly Met Phe Val Tyr Glu  
180 185 190

Pro Asn Leu Ser Thr Tyr His Asn Leu Leu Glu Thr Val Lys Ile Val  
195 200 205

Pro Pro Thr Leu Phe Ala Glu Gln Asp Phe Leu Asn Met Tyr Phe Lys  
210 215 220

Asp Ile Tyr Lys Pro Ile Pro Pro Val Tyr Asn Leu Val Leu Ala Met  
 225 230 235 240

Leu Trp Arg His Pro Glu Asn Ile Glu Leu Asp Gln Val Lys Val Val  
 245 250 255

His Tyr Cys Ala Ala Gly Ala Lys Pro Trp Arg Phe Thr Gly Glu Glu  
 260 265 270

Glu Asn Met Asp Arg Glu Asp Ile Lys Met Leu Val Lys Lys Trp Trp  
 275 280 285

Asp Ile Tyr Asn Asp Glu Ser Leu Asp Tyr Lys Asn Val Val Ile Gly  
 290 295 300

Asp Ser His Lys Lys Gln Gln Thr Leu Gln Gln Phe Ile Glu Ala Leu  
 305 310 315 320

Ser Glu Ala Gly Ala Leu Gln Tyr Val Lys Ala Pro Ser Ala Ala  
 325 330 335

<210> 3

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<212> DNA

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| gattacgtga aaggagtcgt tggtttagcc aaagggttaa ggaaagtcaa atcggcttat | 180 |
| ccactcgtag tagcgatggt acccgacgtc ccggaggaac accgtcgtat acttgtggat | 240 |
| caaggatgca tcgtccgtga aatcgaaccc gtttaccac ccgagaacca aactcagttc  | 300 |
| gccatggctt attacgtcat caactactct aaactccgta tctggaagtt tgtggagtat | 360 |
| agtaaaatga tatatttaga tggagacatt caagtttacg aaaacatcga tcacttgttt | 420 |



|                                                                    |      |
|--------------------------------------------------------------------|------|
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| cacacgccgc aatacaagat cagatattgc caacaatgcc ccgacaaagt ccagtggcca  | 540  |
| aaagcggagc ttggagagcc accggctctt tacttcaacg ccggaatggt cttgtacgag  | 600  |
| cctaacctcg agacttacga ggatctacta cgaacactta aaatcactcc tccgactcct  | 660  |
| ttcgctgaac aggatTTTTT gaacatgtac tTTaagaaaa tctacaagcc gattccttta  | 720  |
| gtgtacaatc tcgtccttgc gatgttatgg cgtcacccag aaaatgtaga gcttggaaaa  | 780  |
| gtcaagggtg ttactactg tgcagcgggt tcgaagccgt ggagatacac agggaaagaa   | 840  |
| gcgaacatgg agaggggaaga tataaaaatg ttagtgaaaa aatggtggga catttacgac | 900  |
| gacgaatcct tggattacaa gaaacctgtt accgttgtgg acacagaggt cgatctcgtg  | 960  |
| aatctgaagc cgttcatcac cgctcttact gaagctggcc ggctcaacta cgtgaccgca  | 1020 |
| ccgtccgctg cttgaatggt gccaggagtt aaaattgtcg gtgg                   | 1064 |

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<210> 5

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caaggatccc ctggcaatca agcagcgga

29

<210> 6

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> synthetic DNA

<400> 6

cgccacagta caagatcggt ta

22

<210> 7

<211> 20

<212> DNA

<213> Artificial Sequence

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<400> 7

catgaagagg cgtatgcagc

20

<210> 8

<211> 20

<212> DNA

<213> Artificial Sequence

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ctttctcggg caagatggca

20

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<211> 20

<212> DNA

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gtgttgacaa gaacctcgct

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<210> 11

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cgcggatccc caccgacaat tttaactcct gg

32

<210> 12

<211> 30

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<223> synthetic DNA

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cgcggatccc tgggtgttgac aagaacctcg etc

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